

Alignment of SDB
w/ Wallach #4

RESULT 4
AAM92795
ID AAM92795 standard; protein; 540 AA.
XX
AC AAM92795
XX
DT 07-MAY-1999 (first entry)
XX
DE Human B1 protein.
XX
KW B1 protein; intracellular mediator; modulator; inflammation; cell death;
KW cell survival pathway; intracellular signalling; AIDS; cancer; human.
XX
OS Homo sapiens.
XX
PN MO9855507-A2
XX
PD 10-DEC-1998
XX
PF 01-JUN-1998; 98WO-IL000255.
XX
XX 05-JUN-1997; 97IL-00121011.
PR 30-JUN-1997; 97IL-00121199.
PR 11-SEP-1997; 97IL-00121746.
XX
PA (YEDA) YEDA RES & DEV CO LTD.
XX
PI Wallach D, Boldin M, Malinin N;
XX

DR MPI; 1999-070258/06.
DR N-PSDB; AAX02558.
XX
XX
PT New B1 protein regulates cell death and cell survival pathways -
PT derivatives, DNA and antibodies, also regulate intracellular inflammation
PT ; for treating AIDS, cancer.
XX
XX
PS Claim 4; Fig 3A; 90pp; English.

XX
XX This invention describes the isolation of a novel human B1 protein which
CC can interact with, intracellular mediators or modulators of inflammation,
CC cell death and/or cell survival pathways, directly or indirectly. Cells
CC can be modulated or mediated in inflammation, cell death or cell survival
CC pathways or another intracellular signalling activity using B1.
CC Conditions such as AIDS and cancer can be treated using B1. Antibodies,
CC oligonucleotides and ribozymes can also be used to regulate the above
CC pathways
XX
XX

SQ Sequence 540 AA;

Query Match 97.8%; Score 227; DB 2; Length 540;
Best Local Similarity 100.0%; Pred. No. 1e-221;
Matches 227; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	6	LSGVSSAIHLCDKKXKELSLNIPVNHGPOESCGSSQLHNSGSPETSRSLPAPQDNDPL	65
Db	314	LSGVSSAIHLCDKKXKELSLNIPVNHGPOESCGSSQLHNSGSPETSRSLPAPQDNDPL	373
QY	66	SRKADPCYFKLHHCNHSWSTISGSORAAFCDHKTTPCSAIIINPLSTAGNSRLQF	125
Db	374	SRKADPCYFKLHHCNHSWSTISGSORAAFCDHKTTPCSAIIINPLSTAGNSRLQF	433
QY	126	GIAQWIOSREDIVNQTEACINQSLDALSLRDLIMKEDYELVSTKPTTSKYRQLDT	185
Db	434	GIAQWIOSREDIVNQTEACINQSLDALSLRDLIMKEDYELVSTKPTTSKYRQLDT	493
QY	186	TDIOGEFAKVIYQKLDNKQKGLQPYPRILVVSRSPSINLQKSM	232
Db	494	TDIOGEFAKVIYQKLDNKQKGLQPYPRILVVSRSPSINLQKSM	540

9201

AAW92795
ID AAW92795 standard; protein; 540 AA.
XX
AC AAW92795;
XX
DT 07-MAY-1999 (first entry)
XX
DE Human B1 protein.
XX
KW B1 protein; intracellular mediator; modulator; inflammation; cell death;
KM cell survival pathway; intracellular signalling. AIDS; cancer; human.
XX
OS Homo sapiens.
XX
PN MO855507-A2.
XX
PD 10-DEC-1998. 1026
XX
PF 01-JUN-1998; 98WO-IL000255.
XX
PR 05-JUN-1997; 97IL-00121011.
PR 30-JUN-1997; 97IL-00121199.
PR 11-SEP-1997; 97IL-00121746.
XX
PA (YEDA) YEDA RES & DEV CO LTD.
XX
PI Wallach D, Boldin N;
XX
DR WPI: 1999-070258/06.
DR N-FSDB; AAX02558.
XX
PT New B1 protein regulates cell death and cell survival pathways -
FT derivatives, DNA and antibodies, also regulate intracellular inflammation
PT ; for treating AIDS, cancer.
XX
PS Claim 4; Fig 3A; 90pp; English.

This invention describes the isolation of a novel human B1 protein which can interact with, intracellular mediators or modulators of inflammation, cell death and/or cell survival pathways, directly or indirectly. Cells can be modulated or mediated in inflammation, cell death or cell survival pathways or another intracellular signalling activity using B1. Conditions such as AIDS and cancer can be treated using B1. Antibodies, oligonucleotides and ribozymes can also be used to regulate the above pathways

Seq. No.:	2,866-237	Length:	540
Score:	227.00	Matches:	227
Percent Similarity:	100.00%	Conservative:	0
Best Local Similarity:	100.00%	Mismatch:	0
Query Match:	42.75%	Indels:	0
DB:	2	Gaps:	0

US-09-771-161A-2 (1-1669) x AA092795 (1-540)	
QY	335 TTGAGAGGTGTTTCAATGGCCATTCCACTTATGTGACAGAGAAATGAAATTTCTCTG 394
Db	314 LeuDimethylserSerAlaIleHsIleucysAspIysIysMetGluSerIeu 333
QY	395 AACATACCTGAAATCATGTGTCCACAGAGAAATCATGTGATCTCTGACCTCCATGAA 454
Db	334 AsnIleProValaAsnHsIleGlyProGlnGluIleuSerIeuSerGlnIleuHsIleGlu 353
QY	455 AATAGGTATCTCTGAACTTCAAGTCCCTGACCTCCTCAAGCAATGATTTT 514
Db	354 AsnSerGlySerProGluIleuSerIeuSerIeuProIleuProGlnAspAspPheIeu 373
QY	515 TCTGAAAGGTCCAGACTGTTATTTATGAGGTGATCACTGCTCTGGAATACAGT 574
Db	374 SerArgIysAlaGlnAspCysTyrPheMetIleuHsIleuIscysProGlnIAsnHsIser 393
QY	575 TGGATAGCACCATTTCTGATCTCAAGAGGCTGATTTCTGTGATCAAGACCACTTCA 634
Db	394 TrpAspSerThrIleSerGlySerGlnArgAlaIlePheCysAspHsIlysthrThrPro 413
QY	635 TGCTCTTCAGCAATATTAATCCATCTGCAAGTCCAGAAATCCAGAACGCTGACGCT 694
Db	414 CysSerSerAlaIleIleAsnProIeuSerThrAlaGlyIleuSerGluArgIleuGlnPro 433
QY	695 GGTATAGCCAGCAGTGGATCCAGAGCAAGCAAGAGACATTTGTGAACCAATGACAGAA 754
Db	434 GlyIleAlaGlnGlnIlePrlIleGlnSerIysArgIleuAspIleValaAsnGlnMetThrGlu 453
QY	755 GCGTGGCTTAAACAGTGGTATGAGCCCTGTGTCAGGACCTGATCATGAAAGAGAGAC 814
Db	454 AlaCysIleuAsnGlnSerIleuAspAlaIleuIleuSerIysAspIleuIleMetGluAsp 473
QY	815 TATGAACTTGTATGACCAAGCTTCAAGAGCACTCAAAAGTCAGACAAATTACTAGACACT 874
Db	474 TyrGlnIleuValaSerThrIlyProThrArgThrSerIysValaArgGlnIleuLeuAspThr 493
QY	875 ACTGACATCCAGAGAGAAATTTGGCCAAAGTTATAGTACAAAAATTTGAAAGATTAACAA 934
Db	494 ThrAspIleGlnGlnGlnGlnPheAlaIlyValaIleValaGlnIlySleuIysAspAsnIys 513
QY	935 CAAATGGGTCTTACGCTTACCCGAAATACCTTGAGGTTTCTATGACCACTTTTAAAT 994
Db	514 GlnMetGlyIleuGlnProIlyProGlnIleuValaIleValaIserArgSerProSerIleuAsn 533
QY	995 TTACTTCAAAATAAAGCATG 1015
Db	534 LeuIleuGlnAsnIlySerMet 540

AA068774	AA068774 standard; protein; 540 AA.
XX	AA068774;
XX	AC
XX	DT
XX	16-MAY-2000 (first entry)
XX	Amino acid sequence of a human phosphorylation effector PHSF-6.
XX	Human; phosphorylation effector; PHSF; proliferative disorder;
KM	immune disorder; neuronal disorder.
OS	Homo sapiens.
XX	Key
XX	Location/Qualifiers

*Admment of Wallach et al. NT
SID 2*

cirrhosis, hepatitis and cancer, developmental disorders e.g. mental retardation, neurological disorders including Alzheimer's disease and Parkinson's disease, autoimmune and inflammatory disorders such as Crohn's disease and diabetes mellitus and finally, viral, bacterial, fungal, parasitic, protozoan or helminthic infections. Furthermore, the polynucleotides encoding KPP may be useful for creating transgenic animals to model human disease, as well as during gene therapy procedures. The current sequence is that of the human KPP cDNA of the invention.

Sequence 1959 BP; 597 A; 430 C; 420 G; 512 T; 0 U; 0 Other;

Very Match

Best Local Similarity 47.0%; Score 785; DB 9; Length 1959;
Matches 785; Conservativity 100.0%; Pred. No. 0;
Mismatches 0; Mismatches 0; Indels 0; Gaps 0;

423 AGGATCATGTGATGCTCTCAGCTCCATGAAATATGATGCTCTCCTGAACTTCAAGGT 482

RESULT 8
ID AAX02558
AC AAX02558; standard: cDNA, 2098 BP.
DT 07-MAY-1999 (first entry)
DE Human B1 cDNA.
KW B1 protein; intracellular mediator; modulator; inflammation; cell survival pathway; intracellular signalling; AIDS; cancer;
OS Homo sapiens.
PN W09855507-A2.
XX

1020

PD 10-DEC-1998.
 XX 01-JUN-1998; 98WO-IL000255.
 XX 05-JUN-1997; 97IL-00121011.
 PR 30-JUN-1997; 97IL-00121199.
 PR 11-SEP-1997; 97IL-00121746.
 XX (YEDA) YEDA RES & DEV CO LTD.
 PI Wallach D, Boldin M, Malinin N;
 XX WPI: 1999-070258/06.
 DR F-PSDB; AAW92795.
 XX New B1 protein regulates cell death and cell survival pathways -
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 CC pathways or another intracellular signalling activity using B1.
 CC Conditions such as AIDS and cancer can be treated using B1. Antibodies,
 CC oligonucleotides and ribozymes can also be used to regulate the above
 CC pathways
 XX
 XX Sequence 2098 BP; 649 A; 452 C; 449 G; 539 T; 0 U; 9 Other;
 XX
 Query Match 46.9%; Score 783; DB 2; Length 2098;
 Best local Similarity 100.0%; Pred. No. 0;
 Matches 783; Conservat 0; Mismatches 0; Indels 0; Gaps 0;
 QY 333 AGTTACAGAGGTTTCAAGTGGCCATTCACCTATGACAGAGAAATGGAATTATCTC 392
 Db 1197 AGTTACAGAGGTTTCAAGTGGCCATTCACCTATGACAGAGAAATGGAATTATCTC 1256
 QY 393 TGAACATACCTGTAAATCATGGTCCACAAGAGAAATCATGTGATCCTTCAGCTCCATG 452
 Db 1257 TGAACATACCTGTAAATCATGGTCCACAAGAGAAATCATGTGATCCTTCAGCTCCATG 1316
 QY 453 AAAATAGTGTCTCTCTGAACTCAAGTCCCTCCAGCTCTCAAGCAATGATTTT 512
 Db 1317 AAAATAGTGTCTCTCTGAACTCAAGTCCCTCCAGCTCTCAAGCAATGATTTT 1376
 QY 513 TATCTAGAAAGCTCAAGCTTTATTTATGAGCTGCATCATGTCCTGGAATCACA 572
 Db 1377 TATCTAGAAAGCTCAAGCTTTATTTATGAGCTGCATCATGTCCTGGAATCACA 1436
 QY 573 GTTGGGATGACACATTTCTGATCTCAAGGCTGCATTTCTGTATCAAGACCATC 632
 Db 1437 GTTGGGATGACACATTTCTGATCTCAAGGCTGCATTTCTGTATCAAGACCATC 1496
 QY 633 CATGCTCTTCAAGCAATTAATCACTCTCACTGAGGAACTCAGAAAGCTTGAGC 692
 Db 1497 CATGCTCTTCAAGCAATTAATCACTCTCACTGAGGAACTCAGAAAGCTTGAGC 1556
 QY 693 CTGGTATAGCCGAGAGTGGATCCAGAGCAAAAGGAGACATTTGAAACCAATGACAG 752
 Db 1557 CTGGTATAGCCGAGAGTGGATCCAGAGCAAAAGGAGACATTTGAAACCAATGACAG 1616
 QY 753 AAGCTGCTTAAACAGTGCCTAGATGCCCTTCTGTCAGAGGACTTGATATGAAGAAG 812
 Db 1617 AAGCTGCTTAAACAGTGCCTAGATGCCCTTCTGTCAGAGGACTTGATATGAAGAAG 1676
 QY 813 ACTATGAACTTTAGTACCAAGCTTCAAGGACCTCAAAAGTCAGCAATTAAGACA 872
 Db 1677 ACTATGAACTTTAGTACCAAGCTTCAAGGACCTCAAAAGTCAGCAATTAAGACA 1736
 QY 873 CTACTGACATCCAGAGAAATTTGCCAAAGTTATAGTACAAAATGAAAGATAC 932

Db 1737 CTACTGACATCCAGAGAAATTTGCCAAAGTTATAGTACAAAATGAAAGATAC 1796
 QY 933 AACAAATGGGCTTACAGCTTACCCGAAATACCTTGATGATCAACATCTTTAA 992
 Db 1797 AACAAATGGGCTTACAGCTTACCCGAAATACCTTGATGATCAACATCTTTAA 1856
 QY 993 ATTACTTCAAAATTAAGCATGTAGTACTGTTTCAAGAAATGTTTCAATA 1052
 Db 1857 ATTACTTCAAAATTAAGCATGTAGTACTGTTTCAAGAAATGTTTCAATA 1916
 QY 1053 AAGGATTTATATCTGTTGCTTTGACTTTTATATAAATCCGTAGATTAAG 1112
 Db 1917 AAGGATTTATATCTGTTGCTTTGACTTTTATATAAATCCGTAGATTAAG 1976
 QY 1113 CTT 1115
 Db 1977 CTT 1979

Alignment for AF027706 w/ NT Wallach

RESULT 2
AA02558
ID AAX02558 standard; cDNA; 2098 BP.
AC AAX02558;
XX
XX 07-MAY-1999 (first entry)
XX
XX Human B1 cDNA.
XX
XX B1 protein; intracellular mediator; modulator; inflammation; cell death;
XX cell survival pathway; intracellular signaling; AIDS; cancer; human; ss.
XX Homo sapiens.
XX MO855507-A2.
XX
XX 10-DEC-1998.
XX
XX 01-JUN-1998; 98MO-IL000255.
XX
XX 05-JUN-1997; 97IL-00121011.
XX 30-JUN-1997; 97IL-00121199.
XX 11-SEP-1997; 97IL-00121746.
XX
XX (YEDA) YEDA RES & DEV CO LTD.
XX
XX Wallach D, Boldin M, Malinin N;
XX WPI: 1999-070258/06.
XX P-PSDB; AAM92795.
XX
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XX cell death and/or cell survival pathways, directly or indirectly. Cells
XX can be modulated or mediated in inflammation, cell death or cell survival
XX pathways or another intracellular signaling activity using B1.
XX Conditions such as AIDS and cancer can be treated using B1. Antibodies,
XX oligonucleotides and ribozymes can also be used to regulate the above
XX pathways
XX
XX Sequence 2098 BP; 649 A; 452 C; 449 G; 539 T; 0 U; 9 Other;
XX
XX Query Match 80.7%; Score 2017.8; DB 3; Length 2098;
XX Best Local Similarity 99.3%; Pred. No. 1.1e-06;
XX Matches 2016; Conservative 9; Mismatches 6; Indels 0; Gaps 0;
XX
XX AF027706
XX GGCACCAAGCTCTAGAAAAGAGTCACTGTTGGAAGAGCAGCGCTGGCTGGGC 60
XX GGCACCAAGCTCTAGAAAAGAGTCACTGTTGGAAGAGCAGCGCTGGCTGGGC 95
XX
XX 61 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 120
XX CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 155
XX 96 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 180
XX CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 215
XX 121 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 240
XX 156 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 275
XX 181 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 300
XX 216 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 330
XX 241 CATTCCGGGAGATGGCGCCCTGTGACCTAGTGTGGCGGCAAAAAGGCTTTCGCGGC 360

276 TCTGACGCGCCCTGCGCACCAATTCCTTACCAAAATCTGCCGACTGCGCTACTGAGCC 335
301 GCGGCGGCTTGGGACCTGTCGTCGCGCCGCGCACGAGACTGGCGGCTGAGTGGCG 360
336 GCGGCGGCTTGGGACCTGTCGTCGCGCCGCGCACGAGACTGGCGGCTGAGTGGCG 395
361 TGAAGCACTGACCATCCACACTGCGCTGCGACAGTGAAGAAGATGCTTTAAGG 420
396 TGAAGCACTGACCATCCACACTGCGCTGCGACAGTGAAGAAGATGCTTTAAGG 455
421 AAGCTGAATTTTACACAAAGCTAGATTAGTACATCTTCCAAATTTGGGAATTTGCA 480
456 AAGCTGAATTTTACACAAAGCTAGATTAGTACATCTTCCAAATTTGGGAATTTGCA 515
481 ATGACCTGAATTTTGGGAATTTAGTACATCTTCCAAATTTGGGAATTTGCA 540
516 ATGACCTGAATTTTGGGAATTTAGTACATCTTCCAAATTTGGGAATTTGCA 575
541 TCTTCAATGAGAAACCTGAATATCTGATGTTGCTGGCAATGGAATTTGCAATCTTGC 600
576 TCTTCAATGAGAAACCTGAATATCTGATGTTGCTGGCAATGGAATTTGCAATCTTGC 635
601 ATGAAATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 660
636 ATGAAATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 695
661 ACTTGAAGCTGAGAAATTTTATGGAATTTGGAATTTGGAATTTGGAATTTGGAATTT 720
696 ACTTGAAGCTGAGAAATTTTATGGAATTTGGAATTTGGAATTTGGAATTTGGAATTT 755
721 GTTATCAAAAGTGGGCGCATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 780
756 GTTATCAAAAGTGGGCGCATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 815
781 AAGGAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 840
816 AAGGAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 875
841 CAGTATCAAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 900
876 CAGTATCAAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 935
901 AAGGAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 960
936 AAGGAGGAGCAATTTATATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 995
961 ATGACCTGTTATATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1020
996 ATGACCTGTTATATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1055
1021 TCTCTCAATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1080
1056 TCTCTCAATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1115
1081 GTTATATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1140
1116 GTTATATGAGAAAGTGGCGCAATTTGGAAGAGTAACTTTCTTGAACCTG 1175
1141 TTATTCAGTAAAGAAAGAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAA 1200
1176 TTATTCAGTAAAGAAAGAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAA 1235
1201 AAGAGAAAGAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAG 1260
1236 AAGAGAAAGAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAG 1295
1261 GTGATCTCTCAAGCTCAATGAAATGAGTTCCTGAAATCTTCAAGGCTCTTGCAG 1320
1296 GTGATCTCTCAAGCTCAATGAAATGAGTTCCTGAAATCTTCAAGGCTCTTGCAG 1355
1321 CTCTCAAGCAATGATTTTATCTGAGAAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAG 1380
1356 CTCTCAAGCAATGATTTTATCTGAGAAAGTCAAGTCAAGTCAAGTCAAGTCAAGTCAAG 1415

1381 ATCACTGCTCTGGAATTCACAGTTGGATAGCACATTTCTGGTTCTCAAAAGGCTGCAT 1440
1416 ATCACTGCTCTGGAATTCACAGTTGGATAGCACATTTCTGGATCTCAAGGCTGCAT 1475
1441 TCTGTATCAAGACCACTCTCATGCTCTTCAAGCAATATAATCACTCTCACTGCAG 1500
1476 TCTGTATCAAGACCACTCTCATGCTCTTCAAGCAATATAATCACTCTCACTGCAG 1535
1501 GAAACTCAGAACGCTCTGACCTGTGTATACCCAGCACTGGATCCAGCAAAAGGAAAG 1560
1536 GAAACTCAGAACGCTCTGACCTGTGTATACCCAGCACTGGATCCAGCAAAAGGAAAG 1595
1561 ACATTTGGAACCAATATGACAGAGCCCTTAAACAGCTAGATGCCCTTCTGTCCA 1620
1596 ACATTTGGAACCAATATGACAGAGCCCTTAAACAGCTAGATGCCCTTCTGTCCA 1655
1621 GGGACTTGATCATGAAGAGGACTATGAACTTTAGTACCAAGCTTCAAGAGACTCCA 1680
1656 GGGACTTGATCATGAAGAGGACTATGAACTTTAGTACCAAGCTTCAAGAGACTCCA 1715
1681 AAGTCAGCAATTACTAGACACTACTGACATCCAAAGGAAATTTGCCAAAGTTATAG 1740
1716 AAGTCAGCAATTACTAGACACTACTGACATCCAAAGGAAATTTGCCAAAGTTATAG 1775
1741 TACAAAATTTGAAGATACAAACAAATGGGCTTCAAGCTTACCCGAAATCTTGTGG 1800
1776 TACAAAATTTGAAGATACAAACAAATGGGCTTCAAGCTTACCCGAAATCTTGTGG 1835
1801 TTTCTAGATCACACCTTTAAATTTACTCTCAAAATTAAGCATGTAGTACTGTTTTTC 1860
1836 TTTCTAGATCACACCTTTAAATTTACTCTCAAAATTAAGCATGTAGTACTGTTTTTC 1895
1861 AAGAAAGAAATGTTTCATTAAGGATATTTATATCTGTTGCTTGACTTTTTTATA 1920
1896 AAGAAAGAAATGTTTCATTAAGGATATTTATATCTGTTGCTTGACTTTTTTATA 1955
1921 TAAATCCGTTGATTAAGCTTTATGAGGTTCTTTGGTAAATATATGTTCTCCTC 1980
1956 TAAATCCGTTGATTAAGCTTTATGAGGTTCTTTGGTAAATATATGTTCTCCTC 2015
1981 CATGACCTGCTGATTTTATTTATTAATTAATCAAGTAAAGTTGAATTTG 2031
2016 CATGACCTGCTGATTTTATTTATTAATTAATCAAGTAAAGTTGAATTTG 2066

RESULT 3

AAZ09246 standard; cDNA, 1931 BP.

ID

AAZ09246;

AC

25-OCT-1999 (first entry)

DT

Human CARD-3 cDNA.

DE

CARD-3; caspase recruitment domain; CARD-4; regulation; detection;
caspase activation; detection; screening; therapy; diagnosis; disease;
apoptotic cell death; Fas/PO-1 receptor complex; TNF receptor complex;
cancer; follicular lymphoma; carcinoma; p53 mutation; viral infection;
hormone-dependent tumour; autoimmune disorder; Alzheimer's disease;
systemic lupus erythematosus; immune-mediated glomerulonephritis; stroke;
Parkinson's disease; amyotrophic lateral sclerosis; retinitis pigmentosa;
spinal muscular dystrophy; cerebellar degeneration; anaemia; drug;
myelodysplastic syndrome; myocardial infarction; cell proliferation;
cell differentiation; cell survival; CARD-4L; CARD-4S; CARD-4Y; CARD-4Z;
human; ds.

KW

Homo. sapiens.

OS

XX

XX

XX

XX

XX

XX

XX

XX

XX

XX

Location/Qualifiers
Key 214..1836
CDS /*tag= a